



IFWO

RAW SEQUENCE LISTING

DATE: 09/02/2004

PATENT APPLICATION: US/10/810,101

TIME: 08:34:27

Input Set : N:\Crf3\RULE60\10810101.raw.txt

Output Set: N:\CRF4\09022004\J810101.raw

1 <110> APPLICANT: Sagner, Gregor
 2 Tabiti, Karim
 3 Gutekunst, Martin
 4 Soong, Richie
 5 <120> TITLE OF INVENTION: Method For Determining The Efficiency Of Nucleic Acid
 Amplifications
 6 <130> FILE REFERENCE: 5443
 7 <140> CURRENT APPLICATION NUMBER: US/10/810,101
 8 <141> CURRENT FILING DATE: 2004-03-25
 9 <150> PRIOR APPLICATION NUMBER: US/09/823,712
 10 <151> PRIOR FILING DATE: 2001-03-30
 11 <150> PRIOR APPLICATION NUMBER: EP/00107036.6
 12 <151> PRIOR FILING DATE: 2000-03-31
 13 <160> NUMBER OF SEQ ID NOS: 8
 14 <170> SOFTWARE: PatentIn Ver. 3.0
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 17 <211> LENGTH: 18
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 19 <213> ORGANISM: Homo sapiens
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 26 <213> ORGANISM: Homo sapiens
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 33 <213> ORGANISM: Homo sapiens
 34 <400> SEQUENCE: 3
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 45 <211> LENGTH: 19
 46 <212> TYPE: DNA
 47 <213> ORGANISM: Homo sapiens
 48 <400> SEQUENCE: 5

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RAW SEQUENCE LISTING ERROR SUMMARY
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY

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